

UNIVERSITI PUTRA MALAYSIA

CHARACTERIZATION OF GRANULOVIRUS AND NUCLEOPOLYHEDROVIRUS ISOLATED FROM SPODOPTERA LITURA

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By
LAU WEI HONG

Thesis Submitted to the School of Graduate Studies, Universiti Putra Malaysia, in Fulfilment of the Requirements for the Degree of Doctor of Philosophy

March 2002



Specially to my husband, my mother, my brothers and sister



Abstract of thesis presented to the Senate of Universiti Putra Malaysia in fulfilment of the requirement for the degree of Doctor of Philosophy

CHARACTERIZATION OF GRANULOVIRUS AND NUCLEOPOLYHEDROVIRUS ISOLATED FROM SPODOPTERA LITURA

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Two baculoviruses were isolated and identified from *Spodoptera litura*; S. litura nucleopolyhedrovirus (SpltNPV) and S. litura granulovirus (SpltGV). The polyhedra of SpltNPV were about 0.9-1.83 µm in diameter containing multiple virions measuring about 100-280 nm wide and 320-410 nm long. The SpltNPV virions contained nucleocapsids (47-60 nm wide and 300-350 nm long) within an envelope, and the size of capsids measured about 58-60 nm wide and 300-330 nm long.

The capsules of SpltGV were about 0.2-0.3 µm wide and 0.45-0.55 µm long containing single virion (60-73 nm wide and 245-267 nm long). The SpltGV nucleocapsids measured approximately 54-60 nm wide and 287-410 nm long, and found singly enclosed within an envelope. The SpltGV capsids measured about 36-58 nm wide and 175-277 nm long.

The restriction endonuclease analyses (REN) revealed that these two baculoviruses did not show any identical restriction pattern. The DNA size of the SpltNPV and the

D P M

SpltGV was estimated to be 132 kb and 124 kb, respectively. The nucleotide sequence analysis of the polyhedrin gene of SpltNPV had 98% sequence identity to the known SpltNPV (accession number: AF037262); while the granulin gene of SpltGV had 81% sequence identity to the granulin gene of Xestia c-nigrum granulovirus (accession number: U70069). Based on the sequence analysis, the SpltNPV and the SpltGV are placed as a taxon of Group II NPV and Group GV, respectively.

Both viruses exhibited general symptoms of polyhedrosis and granulosis. The SpltNPV-infected larvae showed pinkish yellow at the dorsal and lateral sides, while the SpltGV-infected larvae exhibited whitish ventral. The SpltNPV caused a reduction in the larval size while the SpltGV-infected larvae increased in size with bloated integument when lower viral dosages were given. Both viruses infected fat bodies, Malphigian tubules, tracheal matrices, hypodermis, muscles and midguts. The SpltNPV replicated in the nucleus and spread the disease to susceptible tissues within 24-h postinoculation (p.i). The SpltGV was found replicating in both nucleus and cytoplasm, and the disease spread gradually after 48-h p.i. The LD₅₀ of both viruses in neonate larvae of *S. litura* were 9.04x10² polyhedra for SpltNPV and 1.26x10⁴ capsules for SpltGV. The LT₅₀ of both viruses were similar when neonate larvae were fed with similar ranges of viral dosages. The SpltNPV showed a higher virulence in *S. litura* larvae than the SpltGV. The characterization of these baculoviruses is of particular interest in view of its possible use in biological or integrated control.



Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia sebagai memenuhi keperluan untuk ijazah Doktor Falsafah

PENCIRIAN GRANULOVIRUS AND NUCLEOPOLYHEDROVIRUS DARI SPODOPTERA LITURA

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Dua jenis bakulovirus telah dipencilkan and dicirikan dari Spodoptera litura, iaitu S.

litura nukleopolihedrovirus (SpltNPV) dan S. litura granulovirus (SpltGV). Saiz

polibedra SpltNPV adalab lebih kurang 0.9-1.83 µm diameter dan mengandungi

virion berganda yang berukuran 100-280 nm lebar and 320-410 nm panjang. Virion

SpltNPV mengandungi nukleokapsid (47-60 nm lebar dan 300-350 nm panjang)

dalam satu sampul dan kapsid berukuran 58-60 nm lebar dan 300-330 nm panjang.

Kapsul SpltGV adalah lebih kurang 0.2-0.3 μm lebar dan 0.45-0.55 μm panjang dan

mengandungi satu virion (60-73 nm lebar dan 245-267 nm panjang). Saiz

nukleokapsid SpltGV lebih kurang 54-60 nm lebar dan 287-410 nm panjang dan satu

nukleokapsid terdapat terkurung dalam satu sampul. Kapsid SpltGV adalah

berukuran 36-58 nm lebar dan 175-277 nm panjang.

Analysis Pembatasan Endonuklease (REN) menunjukkan kedua-dua bakulovirus tersebut tidak mempunyai corak pembatasan yang sama. Saiz DNA SpltNPV dan SpltGV telah dianggarkan sebesar 132 kb dan 124 kb, masing-masing. Analysis jujukan nukleotida menunjukkan gen polihedrin SpltNPV mempunyai 98% homologi dengan SpltNPV yang dikenali (nombor asesi: AF037262), manakala gen granulin SpltGV mempunyai 81% jujukan sama dengan gen granulin XcGV (nombor asesi: U70069). Berdasarkan analysis jujukan tersebut, SpltNPV dan SpltGV masing-masing diletakkan sebagai satu takson dalam Kumpulan NPV II dan Kumpulan GV.

Kedua-dua virus menghasilkan simtom penyakit polihedrosis dan granulosis yang umum. Larva yang dijangkiti oleh SpltNPV menunjukkan warna kuning kemerahmudaan pada sisi-sisi tepi dan belakang, manakala larva yang dijangkit oleh SpltGV menunjukkan warna putih pada sisi ventral. SpltNPV menyebabkan pengurangan saiz larva, manakala larva yang dijangkiti oleh SpltGV bertambah saiz badan dengan integumen yang mengembang apabila sukatan virus yang rendah diberikan. Kedua-dua virus menjangkiti tisu lemak, tubul Malfigian, matriks trakea, hipodernis, otot dan usus tengah. SpltNPV membiak dalam nukleus dan penyakit merebak ke tisu-tisu yang mudah dijangkiti dalam masa 24 jam selepas jangkitan (p.i.). SpltGV didapati membiak dalam kedua-dua nukleus dan sitoplasma dan penyakit merebak secara perlahan-lahan selepas 48 jam jangkitan (p.i.). LD50 untuk kedua-dua virus dalam larva *S. litura* yang baru lahir adalah 9.04x10² polihedra untuk SpltNPV dan 1.26x10⁴ kapsul untuk SpltGV. LT50 adalah sama untuk kedua-dua virus bila larva yang baru lahir diberi julat sukatan virus yang serupa. SpltNPV menunjukkan kevirulenan yang lebih tinggi daripada SpltGV terhadap larva *S. litura*. Pencirian



kedua-dua baculovirus tersebut adalah diminati khas disebabkan penggunaannya dalam pengawalan biologi atau bersepadu.



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I certify that an Examination Committee met on 11th March 2002 to conduct the final examination of Lau Wei Hong on her Doctor of Philosophy thesis entitled "Characterization of Granulovirus and Nucleopolyhedrovirus Isolated from *Spodoptera litura*" in accordance with Universiti Pertanian Malaysia (Higher Degree) Act 1980 and Universiti Pertanian Malaysia (Higher Degree) Regulations 1981. The Committee recommends that the candidate be awarded the relevant degree. Members of the Examination Committee are as follows:

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DECLARATION

I hereby declare that the thesis is based on my original work except for the quotations and citations which have been duly acknowledged. I also declare that it has not been previously or concurrently submitted for any other degree at UPM or other institutions.

Lau Wei Hong

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LIST OF ABBREVIATIONS

A260 absorption at 260 nm AcfeMNPV Actebia fennica MNPV

AcMNPV Autographa californica MNPV
AcMNPV Autographa californica MNPV
AgMNPV Anticarsia gemmatalis MNPV
AgNPV Anticarsia gemmatalis NPV

AgurSNPV Aglais urticae SNPV AjGV Achaea janata GV

AnfaMNPV Anagrapha falcifera MNPV AnpeNPV Antheraea pernyi NPV

ArceMNPV Archips cerasivoranus MNPV ArceNPV Archips cerasivoranus NPV ArveGV Argyrotaenia velutinana GV BmMNPV Bombyx mori MNPV Bombyx mori NPV

bp base pairs

BusuNPV
BusuSNPV
Busura suppressaria NPV
CfNPV
Choristoneura fumiferana NPV
Choristoneura fumiferana GV
ChmuGV
Choristoneura murinana GV
Choristoneura rosaceana MNPV

CpGV Cydia pomonella GV

CrleGV Cryptophlebia leucotreta GV
DiheSNPV Diprion hercyniae SNPV
DNA deoxyribonucleic acid

dNTP deoxynucleoside triphosphate

EcobSNPV Ectropis obliqua SNPV

EDTA ethylenediamine tetraacetic acid EppoMNPV Epiphyas postvittana MNPV EppoNPV Epiphyas postvittana NPV ErtiSNPV Erannis tiliaria NPV

EsacGV Estigmene acrea GV
EuocGV Euxoa ochrogaster GV
ExapGV Exartema appendiceum GV

GlbiGV Glena bisula GV
GV Granulovirus

h hour

HzSNPV

HabrGV Harrisina brillians GV
HearNPV Helicoverpa armisgera NPV
HycuGV Hyphantria cunea GV
HycuNPV Hyphantria cunea NPV
HzNPV Helicoverpa zea NPV
HzSNPV Helicoverpa zea SNPV

Helicoverpa zea SNPV

UPM

JucoGV Junonia coenia GV kb kilobase pairs LD₅₀ lethal dose 50

LdNPV Lymantria dispar NPV Leucania seperata NPV

LT₅₀ lethal time 50

LymoNPV Lymantria monacha NPV

M micromolar

MaamMNPV Malacosoma americanum MNPV MacoNPV Malacosoma constrictum NPV MadiMNPV Malacosoma disstria MNPV

MARDI Malaysian Agricultural Research and Development Institute

MbMNPV Mamestra brassicae MNPV MbNPV Mamestra brassicae NPV MepeGV Melanchra persicariae GV

min minute

MNPV multiple nucleocapsid polyhedrosis virus

NanaGV Natada nararia GV

Nese SNPV
Nesw SNPV
Nevi SNPV
Nevi SNPV
Neodiprion sertifer SNPV
Neodiprion swainei NPV
Neodiprion virginiana SNPV

nm nanometer

NPV Nucleopolyhedrovirus

nt nucleotide

OpMNPV Orgyia pseudosugata MNPV
OpNPV Orgyia pseudosugata NPV
OranNPV Orgyia anartoides NPV
Orat SNPV Orgyia antiqua SNPV
Orle SNPV Orgyia leucostigma NPV

p.i. post-inoculation

PadoNPV Panaxia dominula NPV
PaflMNPV Panolis flammea NPV
PalaMNPV Pandemis lamprosana NPV

PbGV Pieris brassicae GV\
PCR polymerase chain reaction

PenuNPV Perina nuda NPV

PhopGV Phthorimaea operculella GV PIB Polyhedral Inclusion Body

PiraGV Pieris rapae GV

PlorNPV Plusia orichalcoa NPV
PlscGV Plathypena scabra GV
Probit probability unit

PsunGV Pseudaletia unipuncta GV
PyanGV Pygaera anastomosis GV
RoMNPV Rachiplusia ou MNPV

second

SacaGV Sabulodes caberata GV



SDS sodium dodecyl sulfate
SeMNPV Spodoptera exigua MNPV
SeNPV Spodoptera exigua NPV

SfMNPV Spodoptera frugiperda MNPV SfNPV Spodoptera frugiperda NPV

SNPV single nucleocapsid polyhedrosis virus

SpliMNPV Spodoptera littoralis MNPV SpltNPV Spodoptera litura NPV TAE Tris-acetate-EDTA buffer

Taq Thermos aquaticus

ThliSNPV Thymelicus lineola SNPV
ThorSNPV Thysanoplusia orchalcea SNPV

TipaNPV Tipula paludosa NPV
TnGV Trichoplusia ni GV
TnSNPV Trichoplusia ni SNPV

TrvmSNPV Trichiocampus viminalis SNPV

V volt

v/v volume per volume
w/v weight per volume
WisiNPV Wiseana signata NPV
WisiSNPV Wiseana signata SNPV
XcGV Xestia c-nigrum GV



CHAPTER 1

INTRODUCTION

An insect is considered a pest when its presence causes an economically important loss. Various types of chemicals have been used for controlling pests. They are fast in action and can be used for a broad range of pests. The excessive use of chemical insecticides, however, can result in building up of pest resistance, side effects on beneficial and non-targeted insects, and pollution to the environment that indirectly harm public health. Chemical control has worsened the pest problem.

Integrated pest management (IPM) promotes an alternative to chemical pest control which includes the use of pest-resistant plants, cultural methods, and biological control, and recommends the application of combined methods to minimize the chance of the target insects adapting to any single tactic (Zechendorf, 1995). Biological control utilizes natural living organisms to control a particular pest. These natural enemies can be a predator or parasite (macrobial control), or pathogen (microbial control) (Burges and Hussey, 1971). They normally occur under the conditions of pest outbreaks and are important factors in reducing the density of pest under natural population (Weiser, 1977). These biological control agents neither accumulate in the food chain nor harm the environment, but only make contact with particular targets. According to Debach and Rosen (1991), only 15% of these organisms have been discovered and identified.



The use of microbial control in insect pest is not a new concept. Microbial insecticides mainly refer to bacteria, fungi, viruses, nematodes, protozoa and rickettsiae. *Bacillus thuringiensis* toxin (Bts) is the most successful bioinsecticide and commercially available. However, resistance of pest against the Bt has been reported (Zechendorf, 1995). Fungi, protozoa and nematodes are slow in action and only effective under favorable conditions. Rickettsiae have low specificity to its host and are also pathogenic to warm-blooded animals (Burges and Hussey, 1971). Therefore, viruses are promising biological control agents since they attack the target insects and reprogram the host cells for virus production. Death is the final stage for insect with viral disease.

Baculoviruses are the most commonly and widely studied double-stranded DNA viruses that infect insects. They are divided into *Nucleopolyhedrovirus* (NPV) and *Granulovirus* (GV) based on their morphology. The baculoviruses have been found in over 600 species of arthropods (Blissard and Rohrmann, 1990) and can be as effective as chemical pesticides in controlling specific pests. They are environmentally attractive because they are highly host-specific and, in general, only infect insect species within a limited host range. They have no impact on plants, mammals, fishes, birds or even non-target insects and do not accumulate in food chains. The occluded viruses are very stable and transmitted horizontally among their host. They are ingested by the susceptible larvae, replicate in the host and finally the host dies releasing large quantities of the occlusion bodies into the environment that maximize the chance of other insects to come in contact with the virus and in turn become infected. Vertical transmission, however, occurs through contamination of



female ovipositor during egg laying. The newly hatched larvae will be infected after consuming the virus from the eggshell.

The development of baculoviruses as an ideal IPM candidate is very promising in the near future. Since people are very concerned on the impact of using chemical control, many baculoviruses have been discovered and studied. Miller and Dawes (1978) reported that HzNPV and OpNPV were registered as pesticides by the U.S. Environmental Protection Agency.

In Malaysia, NPV and GV have been found naturally in the diseased larvae of Spodoptera litura. A pathogenicity test of SpltNPV to S. litura larvae was carried out by Sajap et al. (2000) who reported that the larval mortality was dependent on the viral doses. The properties of NPV and GV found in S. litura larvae have not been fully characterized yet. The fundamental studies of these viruses are important to commercial production. Furthermore, these studies are crucial in managing and manipulating the viruses. Since insect virus classification is based on the intrinsic properties of the virus, this thesis concentrates on the basic characterization of both viruses in order to develop a better biopesticide that is environmentally friendly and highly effective for convolling S. litura. Therefore, the objectives to this study are to:

- 1) Isolate and characterise GV and NPV from Spodoptera litura.
- 2) Analyse the DNA of GV and NPV from S. litura.
- 3) Study the pathology of GV and NPV in S. litura.

